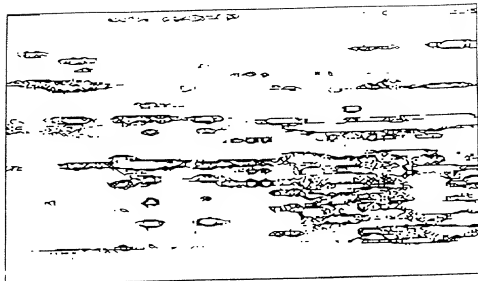
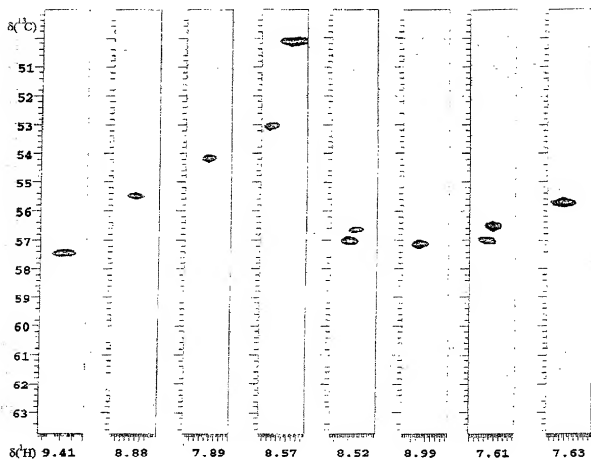


FIGURE 1A



00000000 102201 02000000

FIGURE 1B



# Dipolar Couplings That Depend Only on $\phi(i)$ and $\psi(i)$



Search  $\phi$  and  $\psi$  Until Measured Couplings = Theoretical Couplings

## Packing Secondary Structural Elements

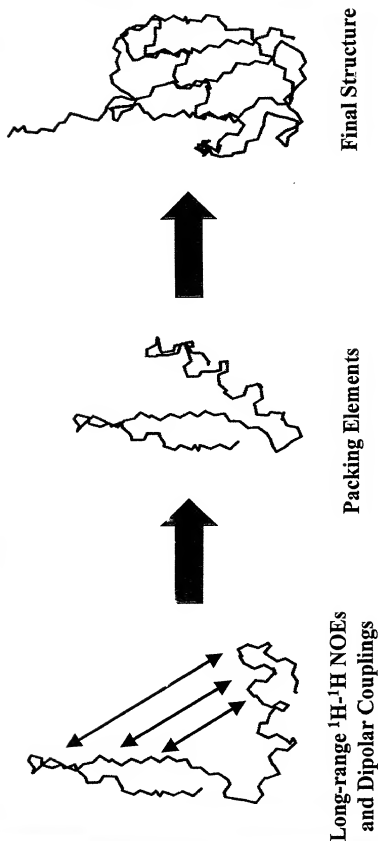
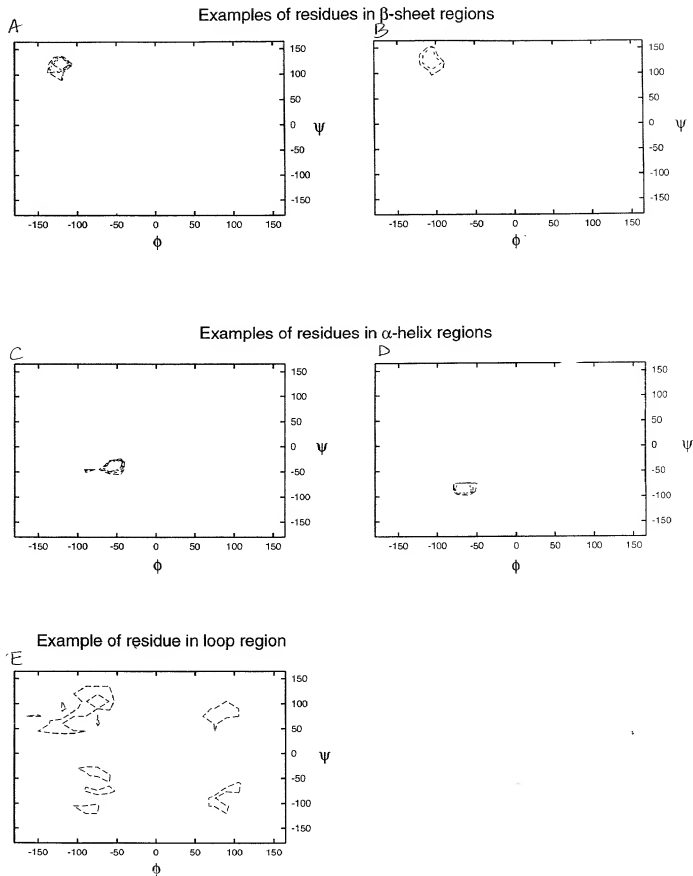


FIGURE 3

FIGURE 4



102201-02028660

0983020.102201

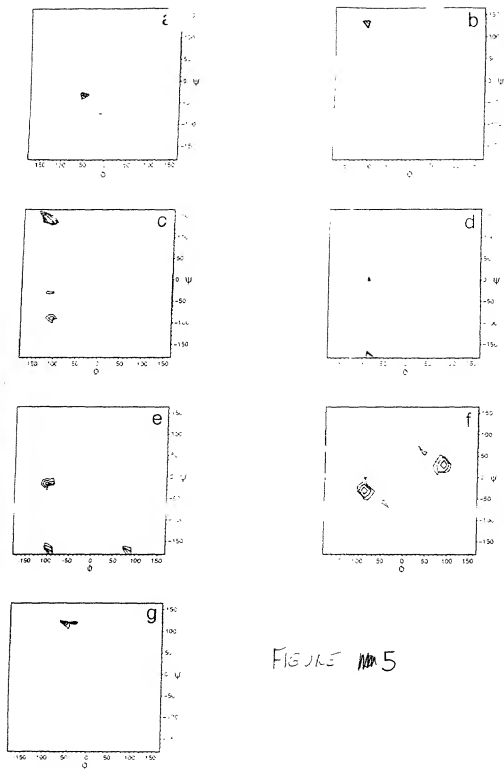


FIGURE 5

Steven W. Homans, et al.  
 RAPID DETERMINATION OF PROTEIN...  
 Atty. Dkt. No. 1496-205  
 Figure 5 of 13



a



b

FIGURE 6

Steven W. Homans, et al.  
RAPID DETERMINATION OF PROTEIN...  
Atty. Dkt. No. 1496-205  
Figure 6 of 13

## FIGURE 7

Generate Linear  
amino-acid chain

Calculate  $\phi, \psi$  angles  
for each peptide pair  
using experimental  
residual dipolar couplings

Fold Linear sequence  
with dihedral angle  
and backbone NOE  
restraints

Refine structure  
using NOE and  
dipolar coupling  
restraints

0963020-102201



# Dipolar Couplings - Powerful Structural Constraints

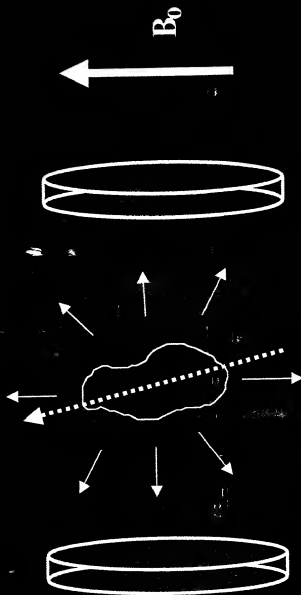


$$D \propto (3 \cos^2 \theta - 1) / r^3$$

$$J + D (\text{Hz})$$



# Measurement of Dipolar Couplings Requires a Weakly Aligned Molecule



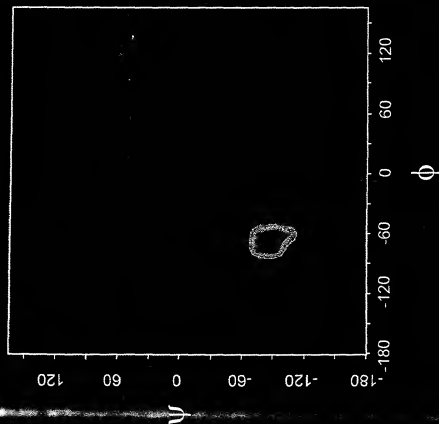
Phospholipid Bicelles



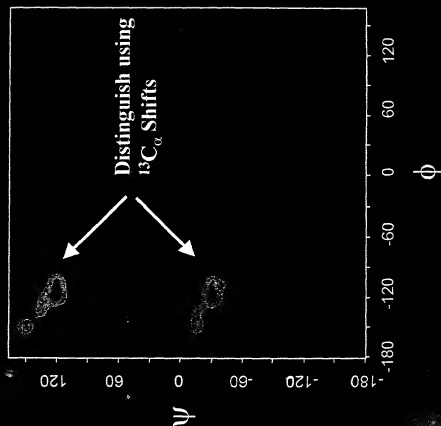
FIGURE 10

# $\phi, \psi$ Mapping Using Residual Dipolar Couplings

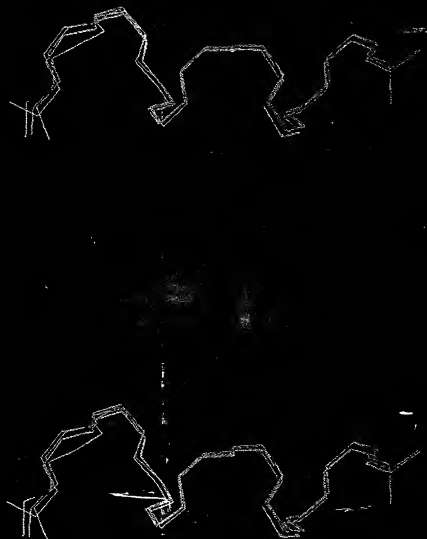
Val 26



Ile 33



# NMR vs. Crystal Structure of $\alpha$ -helix (24-34) Ubiquitin



NMR — Crystal

FIGURE 11

16

# NMR vs. Crystal Structure of $\beta$ -sheet (3-15) Ubiquitin



NMR — Crystal

FIGURE 12

12

# Crystal Structure vs. NMR Global Fold - Ubiquitin

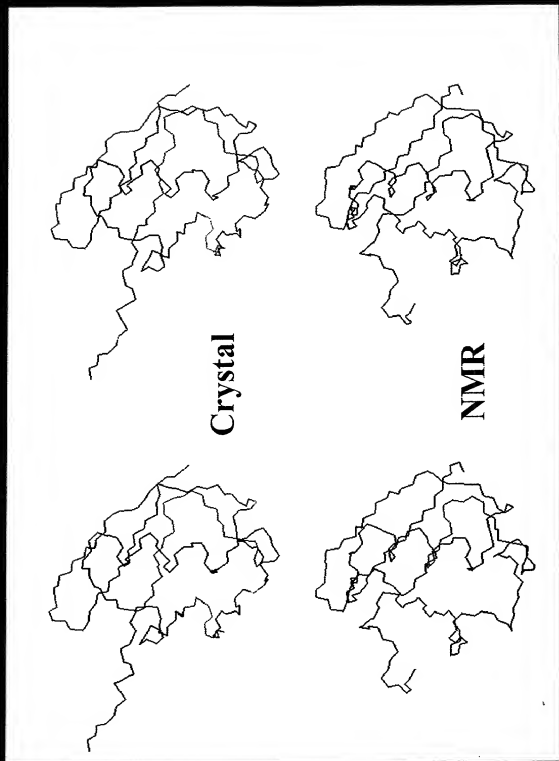


FIGURE 13

9 e n v